

# Marco Terrigno

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1901027/publications.pdf>

Version: 2024-02-01

7  
papers

163  
citations

1684188

5  
h-index

1720034

7  
g-index

11  
all docs

11  
docs citations

11  
times ranked

317  
citing authors

| # | ARTICLE   | IF  | CITATIONS |
|---|---|-----|-----------|
| 1 | A cross-species spatiotemporal proteomic analysis identifies UBE3A-dependent signaling pathways and targets. <i>Molecular Psychiatry</i> , 2022, 27, 2590-2601.                                     | 7.9 | 3         |
| 2 | A eutherian-specific microRNA controls the translation of <i>Satb2</i> in a model of cortical differentiation. <i>Stem Cell Reports</i> , 2021, 16, 1496-1509.                                      | 4.8 | 8         |
| 3 | Pluripotent Stem Cells for Brain Repair: Protocols and Preclinical Applications in Cortical and Hippocampal Pathologies. <i>Frontiers in Neuroscience</i> , 2019, 13, 684.                          | 2.8 | 9         |
| 4 | Neurons Generated by Mouse ESCs with Hippocampal or Cortical Identity Display Distinct Projection Patterns When Co-transplanted in the Adult Brain. <i>Stem Cell Reports</i> , 2018, 10, 1016-1029. | 4.8 | 19        |
| 5 | The microRNA miR-21 Is a Mediator of FGF8 Action on Cortical COUP-TFI Translation. <i>Stem Cell Reports</i> , 2018, 11, 756-769.  | 4.8 | 11        |
| 6 | MicroRNA miR-29 controls a compensatory response to limit neuronal iron accumulation during adult life and aging. <i>BMC Biology</i> , 2017, 15, 9.   | 3.8 | 75        |
| 7 | Assessment of antibody library diversity through next generation sequencing and technical error compensation. <i>PLoS ONE</i> , 2017, 12, e0177574.   | 2.5 | 17        |